



IFWO

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/689,862B

DATE: 07/27/2004

TIME: 11:21:33

Input Set : A:\2004 GENPAX.ST25.txt

Output Set: N:\CRF4\07272004\J689862B.raw

3 <110> APPLICANT: Kim, Sung Hyun  
 5 <120> TITLE OF INVENTION: NOVEL COMPOSITION AND METHODS FOR THE DIAGNOSIS OF LUNG  
 CANCER

7 <130> FILE REFERENCE: GENPAX  
 C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/689,862B

10 &lt;141&gt; CURRENT FILING DATE: 2003-10-20

12 &lt;160&gt; NUMBER OF SEQ ID NOS: 7

14 &lt;170&gt; SOFTWARE: PatentIn version 3.2

16 &lt;210&gt; SEQ ID NO: 1

17 &lt;211&gt; LENGTH: 622

18 &lt;212&gt; TYPE: DNA

19 &lt;213&gt; ORGANISM: Homo sapiens

21 &lt;400&gt; SEQUENCE: 1

22	actacattga	gcatgatgtg	tctcctgaat	gtgtgtttca	tgtgaagtat	tgttctgatt	60
24	aactgacatc	cttgcttaca	agtttcacta	accctttgga	gtttaagcac	aaatgcacaa	120
26	agggaaaaga	ggacgacctg	tttgggggttc	ttttttgcaa	aaacaaacag	tcgcatgctg	180
28	gacgctaaca	ccaagcttac	actgtgtgtg	tgatacggct	gagctgctcc	ataaggctct	240
30	atctttttatc	tgcccaaggc	gtgccctgca	actctggaat	gcagagcagt	tgctgggggtg	300
32	attgacctag	gcacagtggg	gatattttccc	atcttcaaag	ccatgcaaaa	gggcctcctt	360
34	gaccaagaca	caggcctagt	gcttctggaa	tctcaggtta	tcatgtctgg	cctcattgcc	420
36	cctgagacgg	gtgaaaacct	ctctttggag	gagggcgtag	ccagaaacct	cattaatccc	480
38	cagatgtacc	agcagctccg	ggagctacag	gagccctggc	cttaataagc	aggcttactg	540
40	agagcagagg	ccctctttct	gtggtggaag	caattgaaaa	gacaataatc	agtgagacag	600
42	ttggactgaa	aatcttagaa	gt				622

45 &lt;210&gt; SEQ ID NO: 2

46 &lt;211&gt; LENGTH: 268

47 &lt;212&gt; TYPE: DNA

48 &lt;213&gt; ORGANISM: Homo sapiens

50 &lt;400&gt; SEQUENCE: 2

51	gtggtggtgg	gcgctgtaa	tcccagctac	ttggggaggct	gaggcagaga	actgcttgaa	60
53	cccaggaggc	agagggtaca	gtgagccaag	atcgaccac	tgcactccag	cctccagcct	120
55	tcagccttgg	tgacagagca	agactctgtc	tcaaaaagaa	agaaaaagaa	aaagactgtg	180
57	aaagaacaca	catcaaaatg	ttaagcagtg	gtttgtatct	tgaaggacaa	ttttttttta	240
59	ttggaatgtt	tcttctctat	atttttgg				268

62 &lt;210&gt; SEQ ID NO: 3

63 &lt;211&gt; LENGTH: 496

64 &lt;212&gt; TYPE: DNA

65 &lt;213&gt; ORGANISM: Homo sapiens

67 &lt;400&gt; SEQUENCE: 3

68	cgggcccggg	atggactgaa	ccaagaccag	cagccaactt	agaggctcag	ttttaaggcc	60
70	ttgacttggg	atagtaagat	tagagatttc	cagcagtgct	tcctccccgc	acctcccccc	120
72	acccccccgc	ccccgccttt	ttagtgaaga	gaaagtcaca	taaagataac	cattttaaaag	180
74	tgagtaattc	aaggccaggc	gcggtggccc	atgcctgtaa	tcccagcaact	ttgggcggct	240
76	gaggcaggtg	gatcacttga	ggttaggagt	tcgagcccag	cctgggtcaac	atggtgaaac	300

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78 cccgtctcta ctaaaaatat aaaaattagc cgggtgtggt ggcaggcacc tgtaatccca 360
80 gctattaggg aggctgaggc aggagaattg cttgagcctg ggaggcagag gttgcagcga 420
82 gccaaagattg tgccactgta ctccagcctg agcgacggag cgagaatctg tctcaaaaaa 480
84 aaaaaaagat aattca 496
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88 <211> LENGTH: 772
89 <212> TYPE: DNA
90 <213> ORGANISM: Homo sapiens
92 <400> SEQUENCE: 4
93 agttttctaag gatcatgtct gcgagccagg attcccgatc cagagacaat ggccccgatg 60
95 ggatggagcc cgaaggcgctc atcgagggtga gactggagaa atggaattct gtcctcccc 120
97 attacaactt tcagccgtat agagttagag tggcctcttg attgatttcc cagatcatct 180
99 agaagcagct ggtttcccta aagggaggag ggttgtaagc tctgaggctt ttgttagtag 240
101 gcaccagatt ctgtttgctc ggagactaca gctcagctcc accttttcca tgactcaagc 300
103 ttttaatttct ttgcatcccc tagagtaact ggaatgagat tgttgacagc tttgatgaca 360
105 tgaacctctc ggagtcctct ctccgtggca tctacgcta tggttttgag aagccctctg 420
107 ccaccagca gcgagccatt ctaccttgta tcaagggtga gacctctcag tcccagaaga 480
109 cattgtggac tgtccctgac ctgggtagag tggcatctgg ttggtgatgc ccactcata 540
111 tcagccaggg acaaagcaac tccttgttca tcccagcttg gcttttgatc cgtgcccattg 600
113 cctggttcat gccttggaac cataggtttc cttaaagag gtggtattgt agccagctta 660
115 tatttgcata tatagccatg tttctagtcc agcttggtgt gcaatactag atgagttaat 720
117 aactggtcct tgtttctgat ctggttccca ttgtgtaact gtgttgattg gg 772
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121 <211> LENGTH: 705
122 <212> TYPE: DNA
123 <213> ORGANISM: Homo sapiens
125 <400> SEQUENCE: 5
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128 aattgaaacc tggcgcaata gatatagtag cgcaagggaa agatgaaaaa ttatagccaa 120
130 gcataatata gcaaggacta acccctatac cttctgcata atgaattaac tagaaataac 180
132 tttgcaagga gagccaaagc taagaccccc gaaaccagac gagctacctt agaacagcta 240
134 aaagagcaca cccgtctatg tagcaaaaata gtgggaagat ttataggtag aggcgacaaa 300
136 cctaccgagc ctggtgatag ctggttgccc aagatagaat cttagttcaa ctttaaattt 360
138 gccacagaa ccctctaaat ccccttgtaa atttaactgt tagtccaaag aggaacagct 420
140 ctttggacac taggaaaaaa cctttagtag agagtaaaaa atttaacacc catagtaggc 480
142 ctaaaâgacag ccaccaatta agaaagcgtt caagctcaac acccactacc taaaaaatcc 540
144 caaacatata actgaactcc tcacacccaa ttggaccaat ctatcacctt atagaagaac 600
146 taatgttagt ataagtaaca tgaaaacatt ctctccgca taagcctgcg tcagattaaa 660
148 aactgaact gacaattaac agcccaatat ctacatcaac caaca 705
151 <210> SEQ ID NO: 6
152 <211> LENGTH: 301
153 <212> TYPE: DNA
154 <213> ORGANISM: Homo sapiens
156 <400> SEQUENCE: 6
157 tggctcatgg ctacaatccc agcactttgg gaggccgagg caggcagatc accggaggctc 60
159 aggagttcaa gaccagcctg accaacacgg agaaaccccg tcccaactaa aaatacaaaa 120
161 ttagccaggc atggtggcac atgcctgtaa taccagctac tcaggaggct gaggcaggag 180
163 aatgacttga acctgagagg caaatgctgc agtgagccga gatcaggcca ttgcactcca 240
165 gcttgggaaa caagaggcaa aactccgtct ccaaaaaaaaa aaaaaaaaaa aaaaaaaaaa 300

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Input Set : A:\2004 GENPAX.ST25.txt

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167 a 301

170 <210> SEQ ID NO: 7

171 <211> LENGTH: 953

172 <212> TYPE: DNA

173 <213> ORGANISM: Homo sapiens

176 <220> FEATURE:

177 <221> NAME/KEY: misc\_feature

178 <222> LOCATION: (354)..(354)

179 <223> OTHER INFORMATION: n is a, c, g, or t

181 <220> FEATURE:

182 <221> NAME/KEY: misc\_feature

183 <222> LOCATION: (394)..(394)

184 <223> OTHER INFORMATION: n is a, c, g, or t

186 <220> FEATURE:

187 <221> NAME/KEY: misc\_feature

188 <222> LOCATION: (403)..(403)

189 <223> OTHER INFORMATION: n is a, c, g, or t

191 <220> FEATURE:

192 <221> NAME/KEY: misc\_feature

193 <222> LOCATION: (418)..(418)

194 <223> OTHER INFORMATION: n is a, c, g, or t

196 <220> FEATURE:

197 <221> NAME/KEY: misc\_feature

198 <222> LOCATION: (429)..(429)

199 <223> OTHER INFORMATION: n is a, c, g, or t

201 <220> FEATURE:

202 <221> NAME/KEY: misc\_feature

203 <222> LOCATION: (432)..(432)

204 <223> OTHER INFORMATION: n is a, c, g, or t

206 <220> FEATURE:

207 <221> NAME/KEY: misc\_feature

208 <222> LOCATION: (437)..(437)

209 <223> OTHER INFORMATION: n is a, c, g, or t

211 <220> FEATURE:

212 <221> NAME/KEY: misc\_feature

213 <222> LOCATION: (455)..(455)

214 <223> OTHER INFORMATION: n is a, c, g, or t

216 <220> FEATURE:

217 <221> NAME/KEY: misc\_feature

218 <222> LOCATION: (459)..(459)

219 <223> OTHER INFORMATION: n is a, c, g, or t

221 <220> FEATURE:

222 <221> NAME/KEY: misc\_feature

223 <222> LOCATION: (462)..(462)

224 <223> OTHER INFORMATION: n is a, c, g, or t

226 <220> FEATURE:

227 <221> NAME/KEY: misc\_feature

228 <222> LOCATION: (501)..(501)

229 <223> OTHER INFORMATION: n is a, c, g, or t

## RAW SEQUENCE LISTING

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Input Set : A:\2004 GENPAX.ST25.txt

Output Set: N:\CRF4\07272004\J689862B.raw

231 <220> FEATURE:  
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237 <221> NAME/KEY: misc\_feature  
238 <222> LOCATION: (666)..(666)  
239 <223> OTHER INFORMATION: n is a, c, g, or t  
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242 <221> NAME/KEY: misc\_feature  
243 <222> LOCATION: (702)..(702)  
244 <223> OTHER INFORMATION: n is a, c, g, or t  
246 <220> FEATURE:  
247 <221> NAME/KEY: misc\_feature  
248 <222> LOCATION: (709)..(709)  
249 <223> OTHER INFORMATION: n is a, c, g, or t  
251 <220> FEATURE:  
252 <221> NAME/KEY: misc\_feature  
253 <222> LOCATION: (736)..(736)  
254 <223> OTHER INFORMATION: n is a, c, g, or t  
256 <220> FEATURE:  
257 <221> NAME/KEY: misc\_feature  
258 <222> LOCATION: (750)..(750)  
259 <223> OTHER INFORMATION: n is a, c, g, or t  
261 <220> FEATURE:  
262 <221> NAME/KEY: misc\_feature  
263 <222> LOCATION: (785)..(785)  
264 <223> OTHER INFORMATION: n is a, c, g, or t  
266 <220> FEATURE:  
267 <221> NAME/KEY: misc\_feature  
268 <222> LOCATION: (811)..(811)  
269 <223> OTHER INFORMATION: n is a, c, g, or t  
271 <220> FEATURE:  
272 <221> NAME/KEY: misc\_feature  
273 <222> LOCATION: (837)..(837)  
274 <223> OTHER INFORMATION: n is a, c, g, or t  
276 <220> FEATURE:  
277 <221> NAME/KEY: misc\_feature  
278 <222> LOCATION: (844)..(844)  
279 <223> OTHER INFORMATION: n is a, c, g, or t  
281 <220> FEATURE:  
282 <221> NAME/KEY: misc\_feature  
283 <222> LOCATION: (853)..(853)  
284 <223> OTHER INFORMATION: n is a, c, g, or t  
286 <220> FEATURE:  
287 <221> NAME/KEY: misc\_feature  
288 <222> LOCATION: (875)..(875)  
289 <223> OTHER INFORMATION: n is a, c, g, or t  
291 <220> FEATURE:

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Input Set : A:\2004 GENPAX.ST25.txt

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292 <221> NAME/KEY: misc_feature
293 <222> LOCATION: (880)..(880)
294 <223> OTHER INFORMATION: n is a, c, g, or t
296 <220> FEATURE:
297 <221> NAME/KEY: misc_feature
298 <222> LOCATION: (886)..(886)
299 <223> OTHER INFORMATION: n is a, c, g, or t
301 <220> FEATURE:
302 <221> NAME/KEY: misc_feature
303 <222> LOCATION: (896)..(896)
304 <223> OTHER INFORMATION: n is a, c, g, or t
306 <220> FEATURE:
307 <221> NAME/KEY: misc_feature
308 <222> LOCATION: (906)..(907)
309 <223> OTHER INFORMATION: n is a, c, g, or t
311 <220> FEATURE:
312 <221> NAME/KEY: misc_feature
313 <222> LOCATION: (918)..(918)
314 <223> OTHER INFORMATION: n is a, c, g, or t
316 <220> FEATURE:
317 <221> NAME/KEY: misc_feature
318 <222> LOCATION: (925)..(925)
319 <223> OTHER INFORMATION: n is a, c, g, or t
321 <220> FEATURE:
322 <221> NAME/KEY: misc_feature
323 <222> LOCATION: (931)..(931)
324 <223> OTHER INFORMATION: n is a, c, g, or t
326 <220> FEATURE:
327 <221> NAME/KEY: misc_feature
328 <222> LOCATION: (937)..(937)
329 <223> OTHER INFORMATION: n is a, c, g, or t
331 <220> FEATURE:
332 <221> NAME/KEY: misc_feature
333 <222> LOCATION: (940)..(940)
334 <223> OTHER INFORMATION: n is a, c, g, or t
336 <220> FEATURE:
337 <221> NAME/KEY: misc_feature
338 <222> LOCATION: (945)..(945)
339 <223> OTHER INFORMATION: n is a, c, g, or t
341 <220> FEATURE:
342 <221> NAME/KEY: misc_feature
343 <222> LOCATION: (951)..(951)
344 <223> OTHER INFORMATION: n is a, c, g, or t
346 <400> SEQUENCE: 7
347 gatatgaaat gactccctca gacaattttt aaaaagataa gtttttttaa gaccaataaa      60
349 aacctaaggg acaaaaataag acattgatga ttgaaattt ctttgtaaca aatatatacta      120
351 taaatttgaa agcaaaggat agactggaag agaatatattg caatatataa aacaggctaa      180
353 tggtcagtg cacaatatata taacatgctc tcatgtatca attttaaaac aacacccttg      240
355 taaaaaaaaa aaaaaaaaaagg atccaatgag gcagggtaca aataacaaat tcttagcaaa      300

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RAW SEQUENCE LISTING ERROR SUMMARY  
PATENT APPLICATION: US/10/689,862B

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:7; N Pos. 354,394,403,418,429,432,437,455,459,462,501,647,666,702,709  
Seq#:7; N Pos. 736,750,785,811,837,844,853,875,880,886,896,906,907,918,925  
Seq#:7; N Pos. 931,937,940,945,951

## VERIFICATION SUMMARY

DATE: 07/27/2004

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Input Set : A:\2004 GENPAX.ST25.txt

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L:9 M:270 C: Current Application Number differs, Replaced Current Application Number  
L:357 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:300  
L:359 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:360  
L:361 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:420  
L:363 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:480  
L:367 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:600  
L:369 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:660  
L:371 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:720  
L:373 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:780  
L:375 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:840  
L:377 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:900